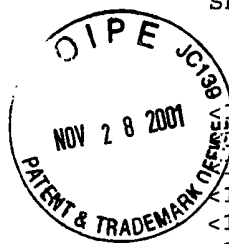


#3

## SEQUENCE LISTING



<110> Smith, Theresa H.  
 <120> PRO-INFLAMMATORY FIBRINOPEPTIDE  
 <130> US 1257/01 (VA)  
 <140> US 09/931,009  
 <141> 2001-08-17  
 <160> 2  
 <210> 1  
 <211> 620  
 <212> PRT  
 <213> Homo sapiens  
 <220>

<400> 1

Ala	Asp	Ser	Gly	Glu	Gly	Asp	Phe	Leu	Ala	Glu	Gly	Gly	Gly	Val	5	10	15
Arg	Gly	Pro	Arg	Val	Val	Glu	Arg	His	Gln	Ser	Ala	Cys	Lys	Asp	20	25	30
Ser	Asp	Trp	Pro	Phe	Cys	Ser	Asp	Glu	Asp	Trp	Asn	Tyr	Lys	Cys	35	40	45
Pro	Ser	Gly	Cys	Arg	Met	Lys	Gly	Leu	Ile	Asp	Glu	Val	Asn	Gln	50	55	60
Asp	Phe	Thr	Asn	Arg	Ile	Asn	Lys	Leu	Lys	Asn	Ser	Leu	Phe	Glu	65	70	75
Tyr	Gln	Lys	Asn	Asn	Lys	Asp	Ser	His	Ser	Leu	Thr	Thr	Asn	Ile	80	85	90
Met	Glu	Ile	Leu	Arg	Gly	Asp	Phe	Ser	Ser	Ala	Asn	Asn	Arg	Asp	95	100	105
Asn	Thr	Tyr	Asn	Arg	Val	Ser	Glu	Asp	Leu	Arg	Ser	Arg	Ile	Glu	110	115	120
Val	Leu	Lys	Arg	Lys	Val	Ile	Glu	Lys	Val	Gln	His	Ile	Gln	Leu	125	130	135
Leu	Gln	Lys	Asn	Val	Arg	Ala	Gln	Leu	Val	Asp	Met	Lys	Arg	Leu	140	145	150
Glu	Val	Asp	Ile	Asp	Ile	Lys	Ile	Arg	Ser	Cys	Arg	Gly	Ser	Cys	155	160	165
Ser	Arg	Ala	Leu	Ala	Arg	Glu	Val	Asp	Leu	Lys	Asp	Tyr	Glu	Asp	170	175	180
Gln	Gln	Lys	Gln	Leu	Glu	Gln	Val	Ile	Ala	Lys	Asp	Leu	Leu	Pro	185	190	195
Ser	Arg	Asp	Arg	Gln	His	Leu	Pro	Leu	Ile	Lys	Met	Lys	Pro	Val	200	205	210

Pro Asp Leu Val	Pro Gly Asn Phe Lys	Ser Gln Leu Gln Lys	Val
215	220		225
Pro Pro Glu Trp	Lys Ala Leu Thr Asp	Met Pro Gln Met Arg	Met
230	235		240
Glu Leu Glu Arg	Pro Gly Gly Asn Glu	Ile Thr Arg Gly Gly	Ser
245	250		255
Thr Ser Tyr Gly	Thr Gly Ser Glu Thr	Glu Ser Pro Arg Asn	Pro
260	265		270
Ser Ser Ala Gly	Ser Trp Asn Ser Gly	Ser Ser Gly Pro Gly	Ser
275	280		285
Thr Gly Asn Arg	Asn Pro Gly Ser Ser	Gly Thr Gly Gly Thr	Ala
290	295		300
Thr Trp Lys Pro	Gly Ser Ser Gly Pro	Gly Ser Thr Gly Ser	Trp
305	310		315
Asn Ser Gly Ser	Ser Gly Thr Gly Ser	Thr Gly Asn Gln Asn	Pro
320	325		330
Gly Ser Pro Arg	Pro Gly Ser Thr Gly	Thr Trp Asn Pro Gly	Ser
335	340		345
Ser Glu Arg Gly	Ser Ala Gly His Trp	Thr Ser Glu Ser Ser	Val
350	355		360
Ser Gly Ser Thr	Gly Gln Trp His Ser	Glu Ser Gly Ser Phe	Arg
365	370		375
Pro Asp Ser Pro	Gly Ser Gly Asn Ala	Arg Pro Asn Asn Pro	Asp
380	385		390
Trp Gly Thr Phe	Glu Glu Val Ser Gly	Asn Val Ser Pro Gly	Thr
395	400		405
Arg Arg Glu Tyr	His Thr Glu Lys Leu	Val Thr Ser Lys Gly	Asp
410	415		420
Lys Glu Leu Arg	Thr Gly Lys Glu Lys	Val Thr Ser Gly Ser	Thr
425	430		435
Thr Thr Thr Arg	Arg Ser Cys Ser Lys	Thr Val Thr Lys Thr	Val
440	445		450
Ile Gly Pro Asp	Gly His Lys Glu Val	Thr Lys Glu Val Val	Thr
455	460		465
Ser Glu Asp Gly	Ser Asp Cys Pro Glu	Ala Met Asp Leu Gly	Thr
470	475		480
Leu Ser Gly Ile	Gly Thr Leu Asp Gly	Phe Arg His Arg His	Pro
485	490		495
Asp Glu Ala Ala	Phe Phe Asp Thr Ala	Ser Thr Gly Lys Thr	Phe
500	505		510

Pro	Gly	Phe	Phe	Ser	Pro	Met	Leu	Gly	Glu	Phe	Val	Ser	Glu	Thr
				515					520					525
Glu	Ser	Arg	Gly	Ser	Glu	Ser	Gly	Ile	Phe	Thr	Asn	Thr	Lys	Glu
				530					535					540
Ser	Ser	Ser	His	His	Pro	Gly	Ile	Ala	Glu	Phe	Pro	Ser	Arg	Gly
				545					550					555
Lys	Ser	Ser	Ser	Tyr	Ser	Lys	Gln	Phe	Thr	Ser	Ser	Thr	Ser	Tyr
				560					565					570
Asn	Arg	Gly	Asp	Ser	Thr	Phe	Glu	Ser	Lys	Ser	Tyr	Lys	Met	Ala
				575					580					585
Asp	Glu	Ala	Gly	Ser	Glu	Ala	Asp	His	Glu	Gly	Thr	His	Ser	Thr
				590					595					600
Lys	Arg	Gly	His	Ala	Lys	Ser	Arg	Pro	Val	Arg	Gly	Ile	His	Thr
				605					610					615
Ser	Pro	Leu	Gly	Lys										
				620										

<210> 2

<211> 4

<212> PRT

<213> Artificial Sequence

<220>

<223> Unknown. Obtained from a commercial source.

<400> 2

Gly Pro Arg Pro